

# Lecture 17. Drift, Ne, Gene Flow

EEB 2245, C. Simon, 28 March 2017

# Last Time..

- Usefulness of Hardy Weinberg
- Early 20<sup>th</sup> century perception of variation in natural populations
- Importance of Lewontin & Hubby 1966
- Importance of genetic variation
- Inbreeding
- Conservation Value
- Intro to Genetic Drift

# This Time..

- Genetic Drift
- Effective population size
- Bottlenecks, Founder Effect
- Gene flow

# Assumptions of Hardy Weinberg

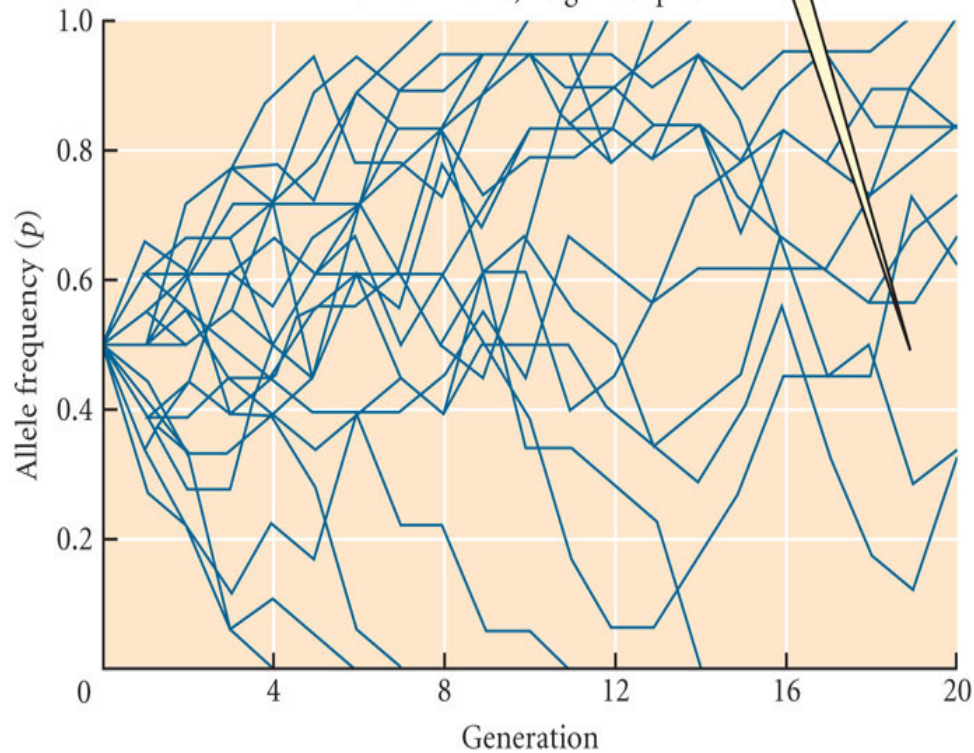
- Diploid population
- No New Mutations
- Random Mating (no inbreeding)
- “infinite” population size (no drift)
- **No Migration of alleles**
- No Natural selection (or tight linkage w another gene under selection)
- Only one population has been sampled.

## Small population

(A)

Oscillations are larger, and alleles are more rapidly fixed or lost, in small populations...

9 individuals, 18 gene copies

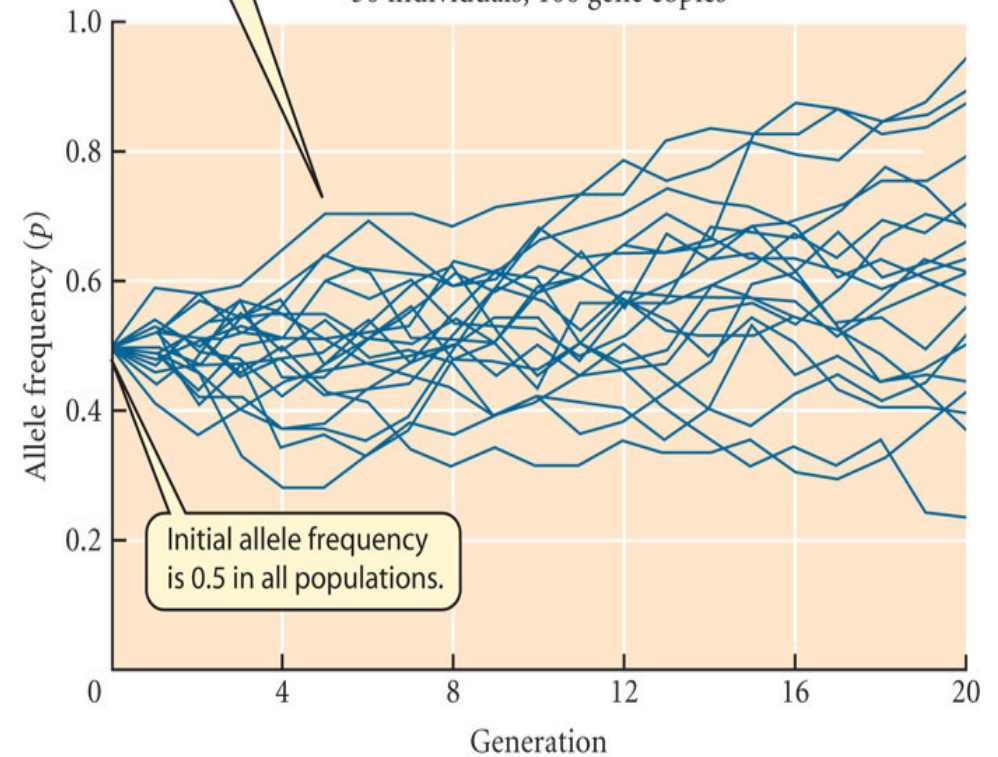


## Large population

(B)

...than in larger ones.

50 individuals, 100 gene copies



Futuyma, 3<sup>rd</sup> Edition, Fig. 10.3

Random genetic drift can result in non-adaptive evolution because one allele is replaced by another (fixed) by chance.

# Genetic Drift

The effect of recessive allele freq. on heterozygosity:

allele freq.:

a =     0.5     0.6     0.7     0.8     0.9     “a” fixed

b =     0.5     0.4     0.3     0.2     0.1     “b” lost

Genotype freq. (at H-W equilibrium):

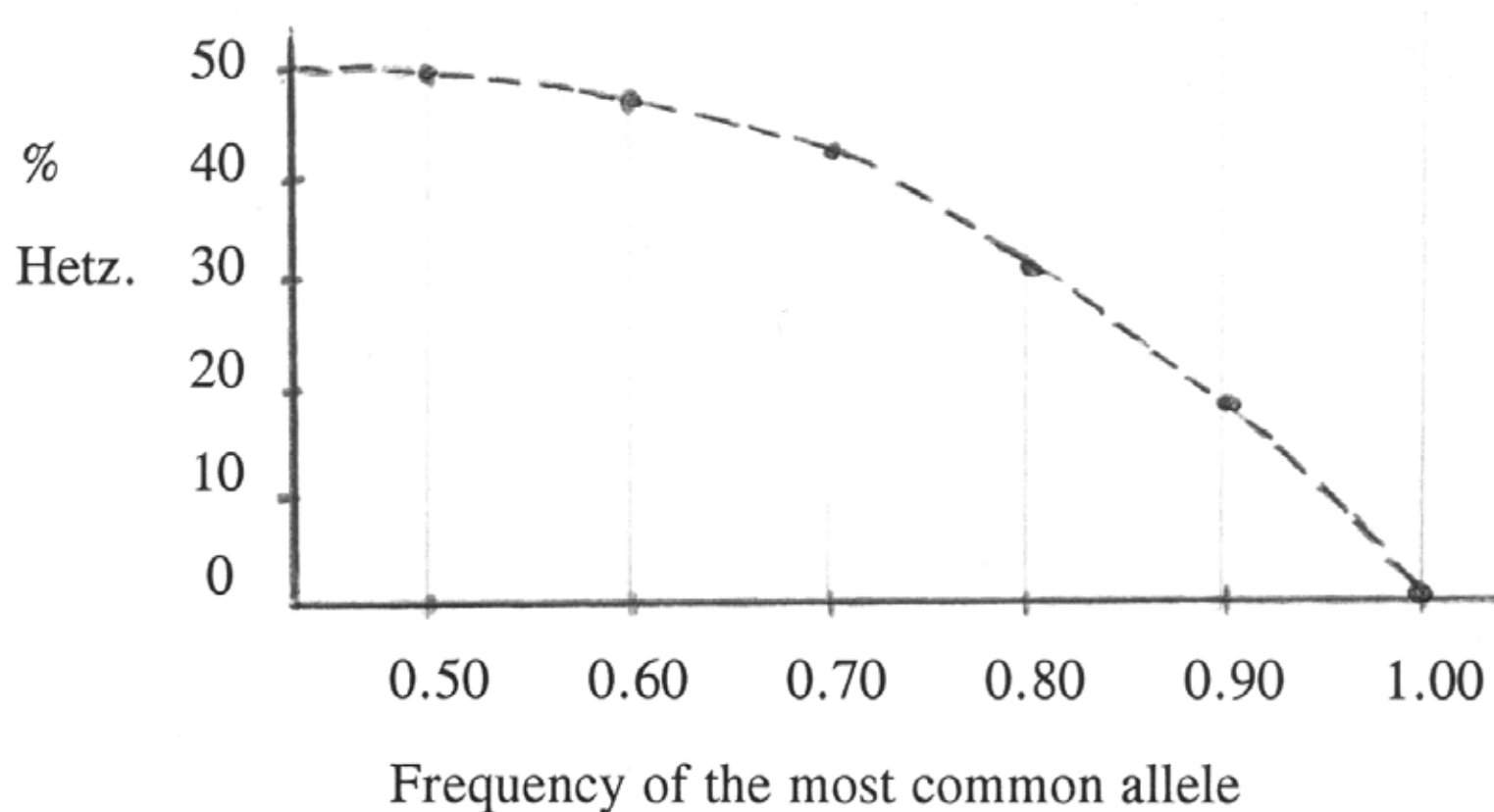
aa =   0.25   0.36   0.49   0.64   0.81

ab =   0.50   0.48   0.42   0.32   0.18

bb =   0.25   0.16   0.09   0.04   0.01

# Genetic Drift

Note that as one allele becomes rare, the proportion of heterozygotes (the heterozygosity of the population) declines as a direct consequence



Effective Population Size,  $N_e$

Reasons for  $N_e$  not equal to census size

# 1) Some males or females don't mate.

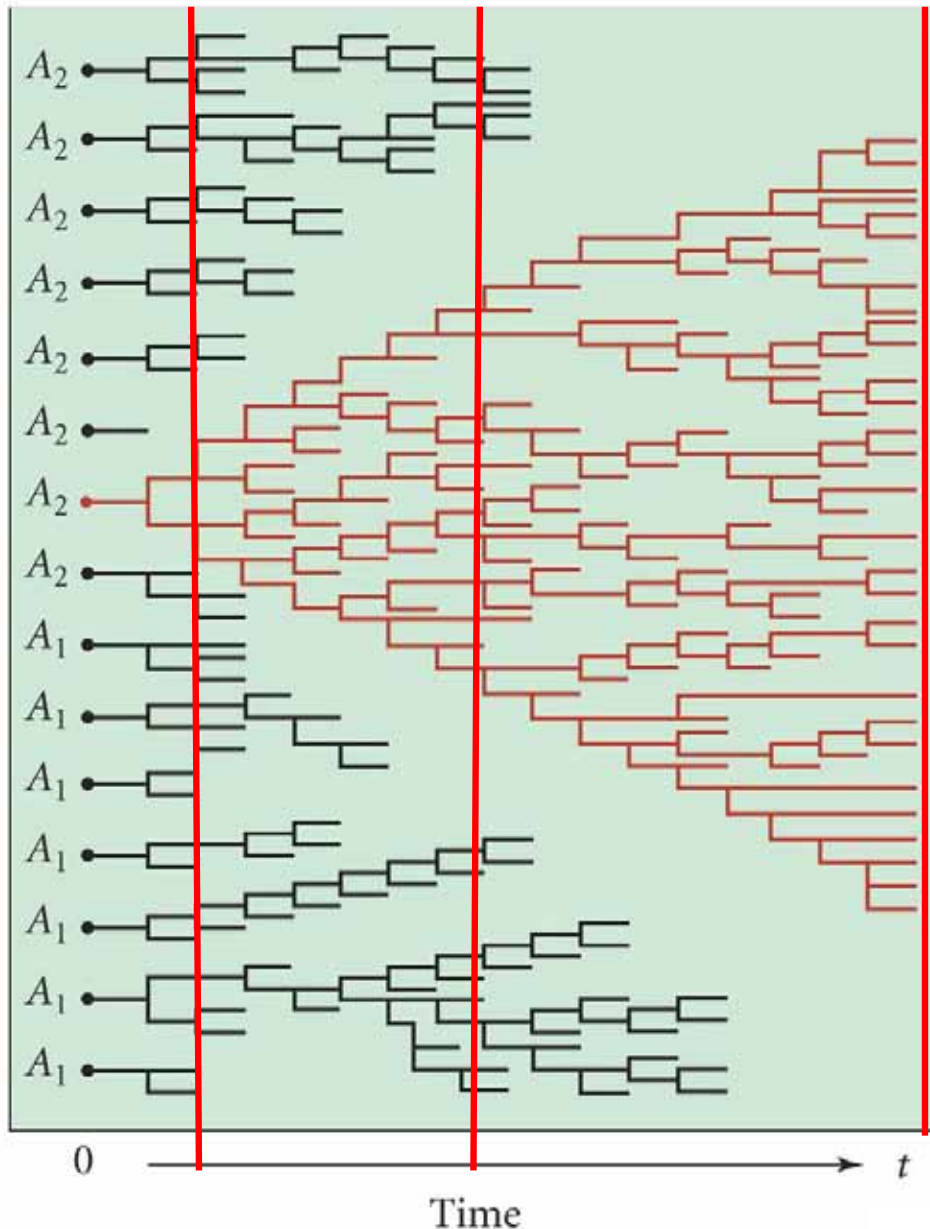
- a) Dominant males, harems (or sex ratio not 1:1)
- b) Sneaky males may achieve secret matings
- c) Observed matings may be thwarted.

Consequence: Some individuals leave fewer progeny than others (increases inbreeding)

Elephant seal territoriality <https://www.youtube.com/watch?v=JrzenOfcmBo>

Video clips: Hedge sparrow from BBC, David Attenborough's "Life of Birds" series, "Finding Partners" <http://www.bbc.co.uk/programmes/p007tx02>

# Coalescence & inbreeding



Initially 15 copies. Most go extinct by chance.

**Average relatedness increases over time**

At time “t” all copies “descend from” or “coalesce to” a single ancestor

**Smaller poplns, faster coalescence**

**Time to coalescence is  $2N$  generations, where  $N$ = popln size**

Additional Reasons for  $N_e$  not equal to census size—

- 2) Sex ratio different from 1 : 1 (similar effect to harems)
- 3) Natural selection causing progeny number to vary among genotypes
- 4) Given overlapping generations, offspring may mate with parental generation
- 5) Fluctuations in population size over time

Drift generally depletes genetic variation

However, in certain circumstances drift can actually enhance genetic variation.

# Applications of genetic drift to conservation Biology

## Conservation Problem:

You are a population biologist and you are called upon to help save an endangered species that if left alone will soon be eliminated in the wild by an introduced predator. The only hope is captive breeding.



Hypothetical organism

**Facts:** Populations are small; genetically isolated--widely scattered. You can only afford to maintain and breed 100 individuals.

**Goal:** Maintain as much genetic variability as possible;  
Keep costs manageable

**Question:** Is it best to collect 100 individuals from one population or 10 individuals from each of ten populations?

# Genetic bottlenecks- small population size

- Causes:
- natural environmental fluctuations (gradual or catastrophic)
- Human intervention- hunting or habitat conversion
- Founder events



<http://animals.nationalgeographic.com/animals/mammals/cheetah/>

# Northern Elephant Seal

- Reduced by hunting
- 20 animals in 1890's.
- Harems- Avg. only 20% of males mate
- 1970's recovered to 30,000
- 1974 Survey: 24 allozymes
- All loci monomorphic (one allele)



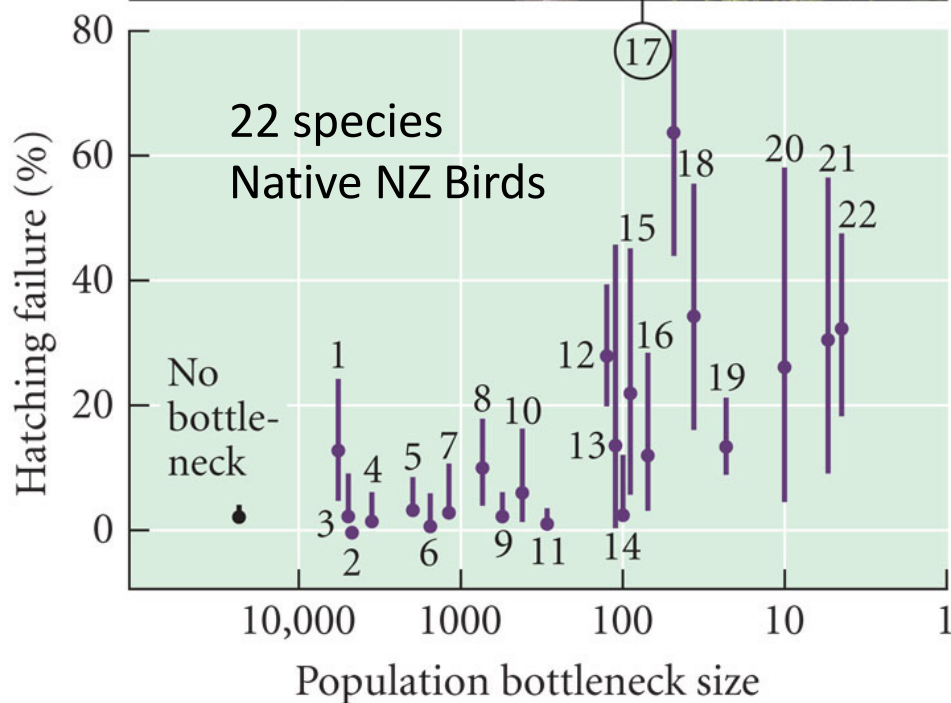
# Failure rate of egg hatching in NZ birds correlated with size of bottleneck

*Strigops habroptila*

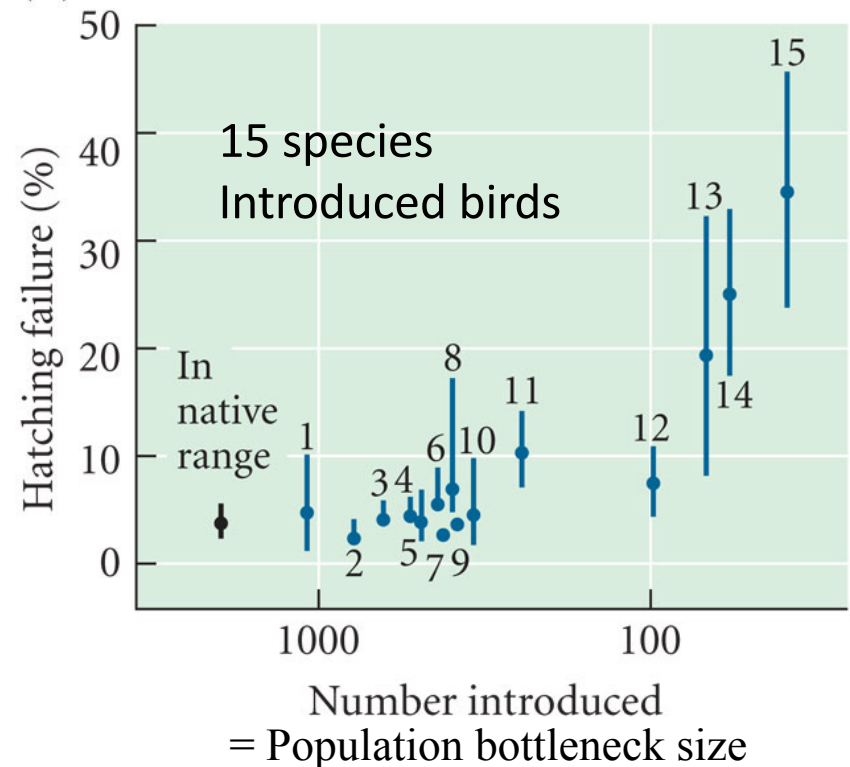


2.3 minute video clip of Kakapo  
narrated by Benedict Cumberbatch  
[https://www.youtube.com/watch?v=E3a88\\_SjJR0](https://www.youtube.com/watch?v=E3a88_SjJR0)

(A)



(B)



Futuyma, 3<sup>rd</sup> Edition, Fig. 10.9

# Finnish People



Genetic Drift in Small Populations  
(original colonists)

Bottleneck- separated for centuries  
(4,000 yr)

Different set of genetic diseases

Pääbo et al. 1996. PNAS

DNA- Y-chromosome & mtDNA

54 Finns

28 Saami (resident Laplanders)

100 Europeans

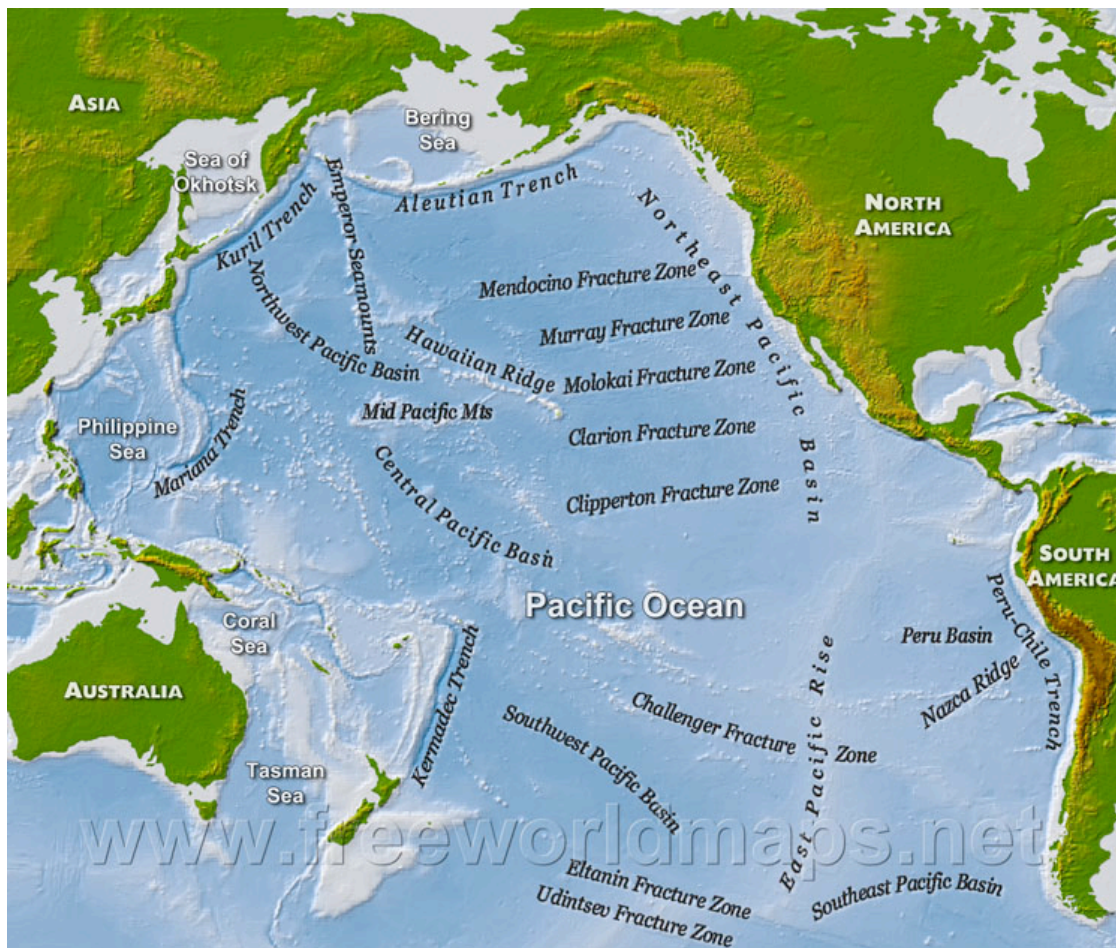
Finns least variable

# Saami People



<http://virtual.finland.fi/finfo/images/people/saami2.jpg>; <http://www.saminuorat.se/media/38957.jpg>; <http://www.arcticpeoples.org/wp-content/uploads/2008/02/ome-photo.jpg>[http://upload.wikimedia.org/wikipedia/commons/thumb/8/87/Saami\\_Family\\_1900.jpg/800px-Saami\\_Family\\_1900.jpg](http://upload.wikimedia.org/wikipedia/commons/thumb/8/87/Saami_Family_1900.jpg/800px-Saami_Family_1900.jpg)

Founder events: one or a few individuals colonize an isolated area



Are oceanic islands deserts of genetic diversity?

# Founder events and bottlenecks

- Two parents (or a pregnant female) founding a new population carry at most four different alleles at a given locus.
- At most loci, one, two or three alleles tend to predominate; founders will most likely carry the most common allele.
- Thus, many alleles will be lost in a bottleneck no matter how long it is
- Extended bottlenecks will tend to reduce variability further due to genetic drift in the small population
- Short bottlenecks will not reduce variability substantially as demonstrated next...

# Effects of founder population size on population heterozygosity:

Given

$H_0$  = Source population proportion of heterozygotes

$N$  = # of colonists

$H_f$  = Heterozygosity of the founder

Population genetics calculate  $H_f = (1 - 1/(2N)) H_0$

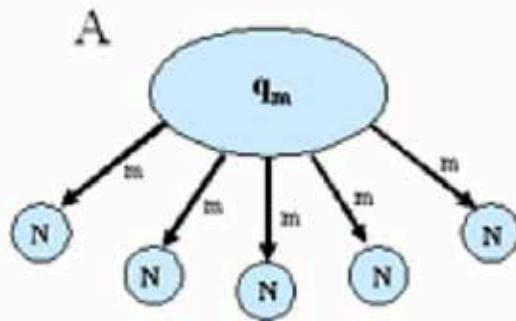
With  $N = 2$  colonists

$$H_f = (1 - 1/(2N)) H_0 \quad \text{or}$$

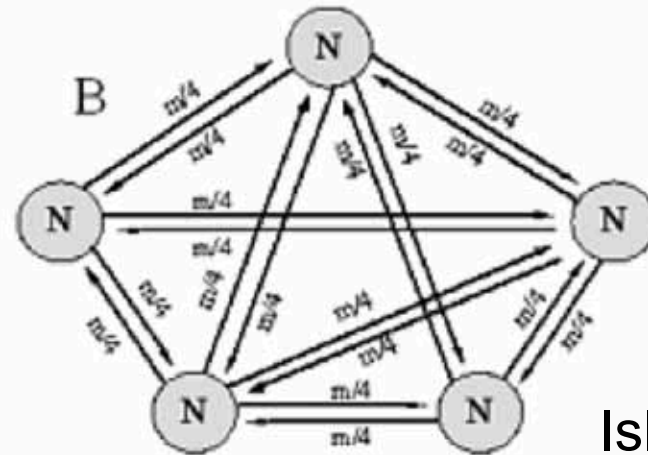
$$H_f = (1 - \frac{1}{4}) H_0 \quad \text{or}$$

$$H_f = \frac{3}{4} H_0$$

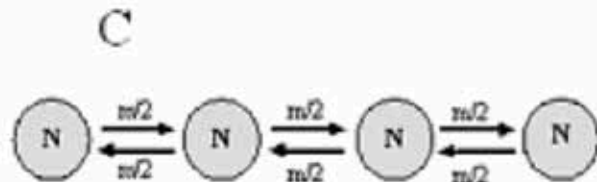
# Models of Gene Flow



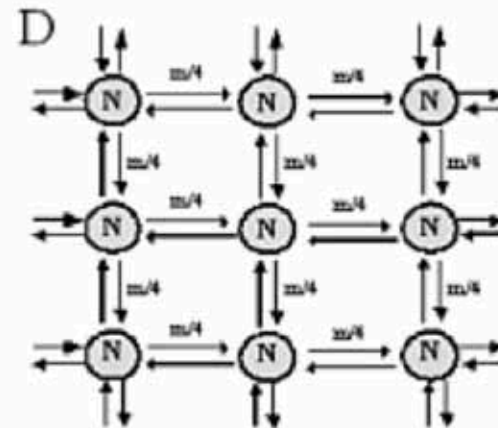
Continent - Island



Island

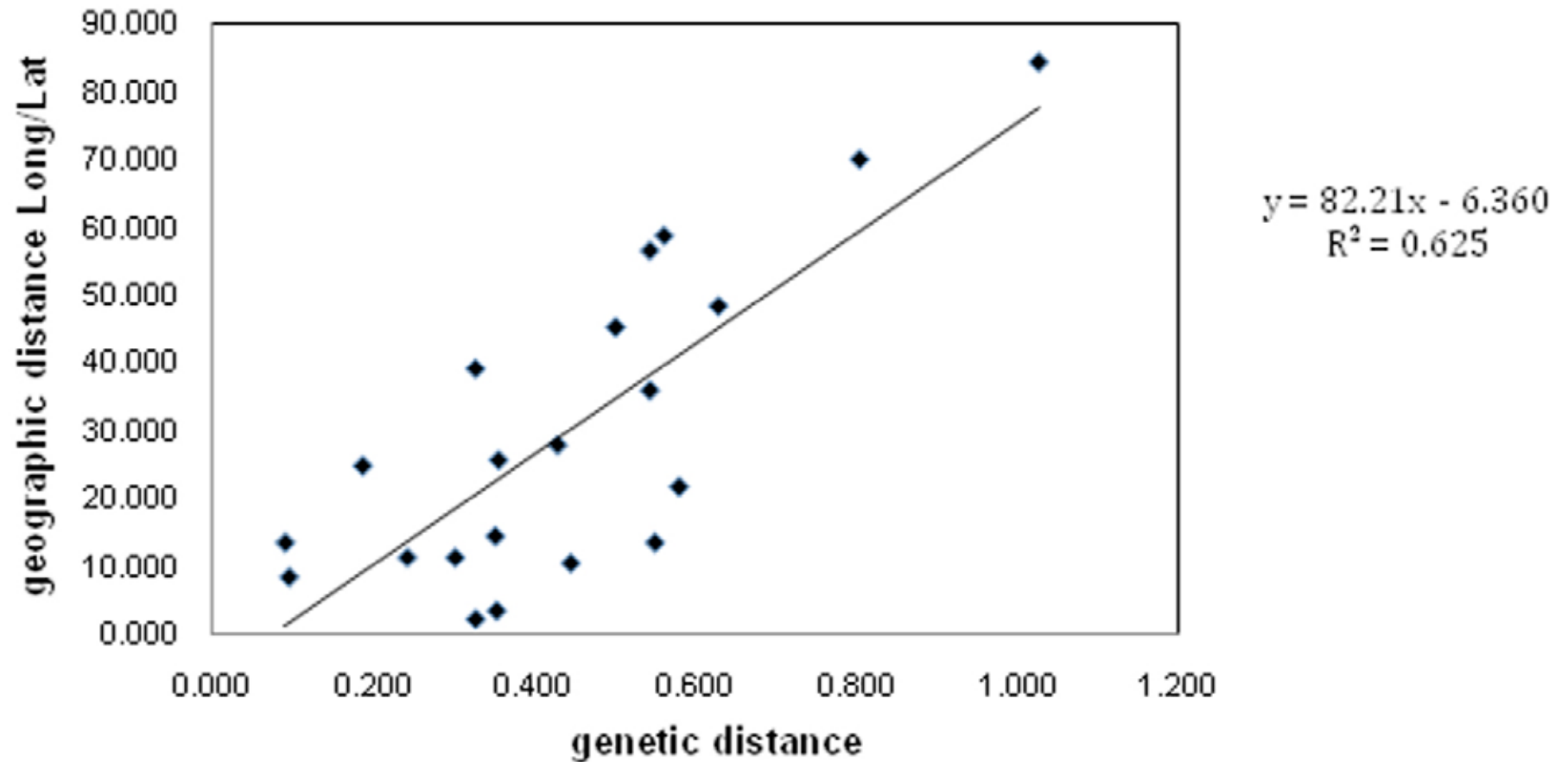


One-dimensional stepping stone



Multi-dimensional stepping stone

# Models of Gene Flow



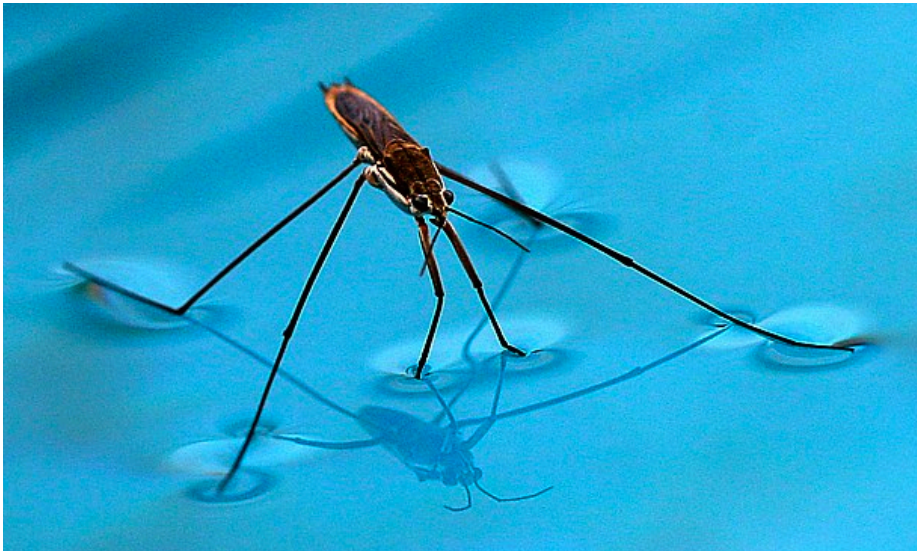
Isolation by distance.

Which model seems most realistic: isolation by distance or perfect panmixia (island model)?

Does migration always = gene flow?

What is the effect of gene flow on among population variation?

# Studies of gene flow in species of high versus low vagility



<http://www.eyefetch.com/image.aspx?ID=452439>



<http://www.encyclopediaofalabama.org/face/Multimedia.jsp?id=m-3167>



The end